

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/937,974

DATE: 10/30/2001

TIME: 14:12:35

Input Set : A:\Dsc64-2.app

Output Set: N:\CRF3\10302001\I937974.raw

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3 <110> APPLICANT: Van Den Boss, Christian
 4 Mbalaviele, Gabriel
 6 <120> TITLE OF INVENTION: Human Mesenchymal DNAs and Expression Products
 8 <130> FILE REFERENCE: 640100-363
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/937,974
 C--> 11 <141> CURRENT FILING DATE: 2001-10-01
 13 <150> PRIOR APPLICATION NUMBER: 60/127,418
 14 <151> PRIOR FILING DATE: 1999-04-01
 16 <150> PRIOR APPLICATION NUMBER: 60/148,800
 17 <151> PRIOR FILING DATE: 1999-08-13
 19 <160> NUMBER OF SEQ ID NOS: 29
 21 <170> SOFTWARE: PatentIn Ver. 2.1
 23 <210> SEQ ID NO: 1
 24 <211> LENGTH: 1974
 25 <212> TYPE: DNA
 26 <213> ORGANISM: Artificial Sequence
 28 <220> FEATURE:
 29 <223> OTHER INFORMATION: Description of Artificial Sequence:cDNA derived
 30 from human mesenchymal stem cells after treatment
 31 with growth supplement
 33 <400> SEQUENCE: 1
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 36 tgttcattggc aacacgagaa cttctgcgga tcccccaagc agccctggcc aagccaatct 180
 37 ctataacctac aaacctagtg tccctctttt ctgcctatgt tgaccggcag aaactgaact 240
 38 tgcctggagac aaaactgcag ctagtctcagg ggatacgata aaagatctcc aaatgtgtcc 300
 39 tgtacctcct tttggctgcc acctgcactg ctgccatcac caatggagtg tttttaatga 360
 40 ggggaaggaa gtagcttttt ccccaagca aagtcttgtg ggatcgatc ctgtttacag 420
 41 ggggttgtctc tctaaatgtc agataattcc ccactgctct atgaaatttg gctgggtgat 480
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 43 tctgttttgc attttctttg cctagagac acaagtgtaa tctctccctt tatccctcca 600
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 46 gatatacaca cacacataga tacacacaca cacaccccat acatgtattt atgtggtctt 780
 47 cagagggtcc ttaaagaatg aatttttagat tgaaaaatat ttagtgtctt cattacctct 840
 48 tctaaacaca aaccagctga tgtattttaa tctgtttctg ttctatcttg taattaat 900
 49 ggtgggttct acttgtttta acataataa agagtatgca gcacgtttta taaaatcaga 960
 50 actttaattt ggttatgcgc caggcttagg ctgagaagtc ctttttcttc ttccacactt 1020
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 58 cccactcagc aatcaaccag actaaggagt tttgatccct agtgattaca gccctgaaga 1500
 59 aaattaaatc tgaattaat ttacatggcc ttctgtatct tctgtctgtt cttacttttt 1560

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60 cgaatgtagt tggggggtgg gagggacagg ttatggtatt taaagagaat aaacattttg 1620
61 cactatcatg tattgtacaa cagtaagatc ctctgtttaa accagctgtc ctgttctcca 1680
62 tctccatttc tcccatgct gtaaccccaq gctccaccag ctgttcccca gtgatgttac 1740
63 ctacttctcc tctacattg tctactgacc atttccacta catcccttcc ctaccttccc 1800
64 ttccaaacca atcaaaqtga tacttgatta ttactctctc ctactgtgac ttatctcttt 1860
65 ttgtttgcat tggttctaat taatgaaaat aaaagtctct aaatttacct ttatatagg 1920
66 tattgtaaat aaaaacaaat tgtatactta aaaaaaaaaa aaaaaaaaaa aaaa 1974

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69 <210> SEQ ID NO: 2

70 <211> LENGTH: 410

71 <212> TYPE: PRF

72 <213> ORGANISM: Artificial Sequence

74 <220> FEATURE:

75 <223> OTHER INFORMATION: Description of Artificial Sequence: Putative
 76 protein derived from ORF cDNA of SEQ ID NO:1

78 <400> SEQUENCE: 2

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83 20 25 30
85 Pro Val Gly Phe Cys Leu Leu Val Leu Arg Leu Phe Leu Gly Ile His
86 35 40 45
88 Val Phe Leu Val Ser Cys Ala Leu Pro Asp Ser Val Leu Arg Arg Phe
89 50 55 60
91 Val Val Arg Thr Met Cys Ala Val Leu Gly Leu Val Ala Arg Gln Glu
92 65 70 75 80
94 Asp Ser Gly Leu Arg Asp His Ser Val Arg Val Leu Ile Ser Asn His
95 85 90 95
97 Val Thr Pro Phe Asp His Asn Ile Val Asn Leu Leu Thr Thr Cys Ser
98 100 105 110
100 Thr Pro Leu Leu Asn Ser Pro Pro Ser Phe Val Cys Trp Ser Arg Gly
101 115 120 125
103 Phe Met Glu Met Asn Gly Arg Gly Glu Leu Val Glu Ser Leu Lys Arg
104 130 135 140
106 Phe Cys Ala Ser Thr Arg Leu Pro Pro Thr Pro Leu Leu Leu Phe Pro
107 145 150 155 160
109 Glu Glu Glu Ala Thr Asn Gly Arg Glu Gly Leu Leu Arg Phe Ser Ser
110 165 170 175
112 Trp Pro Phe Ser Ile Gln Asp Val Val Gln Pro Leu Thr Leu Gln Val
113 180 185 190
115 Gln Arg Pro Leu Val Ser Val Thr Val Ser Asp Ala Ser Trp Val Ser
116 195 200 205
118 Glu Leu Leu Trp Ser Leu Phe Val Pro Phe Thr Val Tyr Gln Val Arg
119 210 215 220
121 Trp Leu Arg Pro Val His Arg Gln Leu Gly Glu Ala Asn Glu Glu Phe
122 225 230 235 240
124 Ala Leu Arg Val Gln Gln Leu Val Ala Lys Glu Leu Gly Gln Thr Gly
125 245 250 255
127 Thr Arg Leu Thr Pro Ala Asp Lys Ala Glu His Met Lys Arg Gln Arg
128 260 265 270

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130 His Pro Arg Leu Arg Pro Gln Ser Ala Gln Ser Ser Phe Pro Pro Ser
131          275          280          285
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134          290          295          300
136 Lys Glu Val Leu Pro His Val Pro Leu Gly Val Ile Gln Arg Asp Leu
137 305          310          315          320
139 Ala Lys Thr Gly Cys Val Asp Leu Thr Ile Thr Asn Leu Leu Glu Gly
140          325          330          335
142 Ala Val Ala Phe Met Pro Glu Asp Ile Thr Lys Gly Thr Gln Ser Leu
143          340          345          350
145 Pro Thr Ala Ser Ala Ser Lys Phe Pro Ser Ser Gly Pro Val Thr Pro
146          355          360          365
148 Gln Pro Thr Ala Leu Thr Phe Ala Lys Ser Ser Trp Ala Arg Gln Glu
149          370          375          380
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152 385          390          395          400
154 Phe Thr Glu Arg Arg Ala Gln Glu Ala Asp
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159 <211> LENGTH: 1619
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161 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA derived
165 from human mesenchymal stem cells after treatment
166 with growth supplement
168 <400> SEQUENCE: 3
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170 aagaatccag gcccttccac gcgcgtgtgg gtgcgggggc cccgaagtgc tcgtggttcc 120
171 ccgctaggtc tccgctgggg caggaaccgg aatcatgggt gggaccacca gcaccgcgcg 180
172 ggtcaccttc gaggcggacg agaatgagaa catcacctgt gtgaagggca tccggttttc 240
173 ggaatatgtg attgatcgaa tgaaggaatc ctctccatct ggttcgaagt ctacgcggta 300
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175 gctggcattg gagcaagcca agaagaatc cgaagatcag aaacgactaa agcaagccaa 420
176 agagctggac cgagagaggg ctgctgccaa tgagcagtta accagagcca tcttccggga 480
177 gaggatatgt agcagaggag aacgcgctaa ggcacaaagc ctggctaggg agctggaaga 540
178 gaaagaccga gtgctaaaga agcaggatgc attctacaaa gaacagctgg ctgactgga 600
179 ggagaggagc tcagagtctc acagagtcac cactgaacaa tatcagaaag ctgctgaaga 660
180 gtggaagcaa agttcaagcg atatgagtct catccagtct gtgctgatct gcaggccaaa 720
181 attcttcagt gttaccgtga gaacacccac cacagccctca aatgctccgc tctggccacc 780
182 cagtatatgc actgtgtcaa tcatgccaaa cagagcatgc ttgagaaggg aggataaaaa 840
183 ctttcagaat gagcaaaaca ccataacagt taattccaga gatggaacat ttttttctct 900
184 agtgagaaaa caaccattt gaagagaaga ccactaatga gaagaccact aaagagagac 960
185 atcaagaagt gattcagcag aatcatttca cgttttgaac acgacagatt tgaaggggcca 1020
186 aagccttgat cagggatcag tcattaaagg acactcttga gtattagtaa accctcttat 1080
187 gatgattaaa agagaagggc agccctctcc accttttggt actttctatt caacttgcac 1140
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191 agtgcctggaat ggaattgtct aagctgtggc atgttggttt tctctttctt ttctcctttg 1380
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193 aaggagagaa aatgtacctc ttctactgga ataattttta tgattacaag tgaqataaag 1500
194 tatttttate aatatgaagg caaccttggc tgataaaacc tctatagtga atactcacat 1560
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201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: Description of Artificial Sequence:Putative
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212     20             25             30
214 Ile Asp Arg Met Lys Glu Ser Ser Pro Ser Gly Ser Lys Ser Gln Arg
215     35             40             45
217 Tyr Ser Gly Ala Tyr Gly Ala Ser Val Ser Asp Glu Glu Leu Lys Arg
218     50             55             60
220 Arg Val Ala Glu Glu Leu Ala Leu Glu Gln Ala Lys Lys Glu Ser Glu
221     65             70             75             80
223 Asp Gln Lys Arg Leu Lys Gln Ala Lys Glu Leu Asp Arg Glu Arg Ala
224     85             90             95
226 Ala Ala Asn Glu Gln Leu Thr Arg Ala Ile Leu Arg Glu Arg Ile Cys
227     100            105            110
229 Ser Glu Glu Glu Arg Ala Lys Ala Lys His Leu Ala Arg Gln Leu Glu
230     115            120            125
232 Glu Lys Asp Arg Val Leu Lys Lys Gln Asp Ala Phe Tyr Lys Glu Gln
233     130            135            140
235 Leu Ala Arg Leu Glu Glu Arg Ser Ser Glu Phe Tyr Arg Val Thr Thr
236     145            150            155            160
238 Glu Gln Tyr Gln Lys Ala Ala Glu Glu Val Glu Ala Lys Phe Lys Arg
239     165            170            175
241 Tyr Glu Ser His Pro Val Cys Ala Asp Leu Gln Ala Lys Ile Leu Gln
242     180            185            190
244 Cys Tyr Arg Glu Asn Thr His Gln Thr Leu Lys Cys Ser Ala Leu Ala
245     195            200            205
247 Thr Gln Tyr Met His Cys Val Asn His Ala Lys Gln Ser Met Leu Glu
248     210            215            220
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255 <211> LENGTH: 1917
256 <212> TYPE: DNA
257 <213> ORGANISM: Artificial Sequence
259 <220> FEATURE:
260 <223> OTHER INFORMATION: Description of Artificial Sequence:cDNA derived

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Input Set : A:\Dsc64-2.app

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261      from human mesenchymal stem cells after treatment
262      with growth supplement
263 <400> SEQUENCE: 5
265 ggcacgagga gagaqqtgtt gaaactcaag atcccccatqg acaacaagga ggtccccggc 60
266 gaggcgcccg cgcctccgcg cgaaccggcg cgtccccacg cgtccccca ctcggcgccg 120
267 gctcttcgcg ggcgtccac gctggcgag cgcgcgcga cgcacacgg cgaacggccc 180
268 ttccggtgca ccgagtgccg gcggcgcttc tcacagaagt cggcgctgac caaacacggc 240
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272 ggcgagaaag cgtacgcgtg ccgggaactc ggaacgcctt ttgcggcgag ctgctgcctg 480
273 gcgcgcacc gacgcagca caggggcgag cggcctacg cttgcgcga ctcggcgacg 540
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275 caccgctgcg ctgtgtgtgg ccgtcgcttc ggcacacgct ccaacctggc ggaacacggc 660
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294 aaacaacaa accaaaccaa acaaaaaaaa tctcaaaagg attggacctc gcagctcatg 1800
295 cctgtaattc ccagcacttt gggaggcgga ggcaggagga tctcttgaag tcaagagttt 1860
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300 <211> LENGTH: 275
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302 <213> ORGANISM: Artificial Sequence
303 <220> FEATURE:
304 <223> OTHER INFORMATION: Description of Artificial Sequence: Putative
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312 Pro Ala Arg Pro His Ala Cys Pro Asp Cys Gly Arg Ala Phe Ala Arg
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315 Arg Ser Thr Leu Ala Lys His Ala Arg Thr His Thr Gly Glu Arg Pro
316 35 40 45

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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date